

Figure 1

CTCAAGTGTATACGTCACGACACTGGCCGTCCGCTCAGGACTGGAACTAGCCGGCTCGCTCAAACCCCCGGGCTTTACATTAGGAGTCAGTGG	100
GGGAGAGTCCTAGGATTTCTAGTCMAACTGACAGCCCTGGACTTTGGACCTTCGTGAAGTCCTCTGCTGGAAAGCTGAGACTTCATGCC ATG	199
M 1	
GAA CAC CCC CTC TTT GGC TGC CTG CGC ACC CCC CAC CCC ACA GCG CAA GCG TTG CAC CCC TTC TCG CAG TCT TCT	274
E H P L F G C L R S P H A T A Q G L H P F S Q S S	26
CTG CCC CTC CAT CGA AGA TCT GAC CAC ATG TCC TAC CCC GAA CTC TCC ACA TCT TCC TCG TCT TCC ATA ATC CGG	149
L A L H G R S D H M S Y P E L S T S S S S C I I A	51
CGA TAC CCC ATT GAG GAG CGC ATG TTT GCC AGC CAG CAT CAC ACC CGG CAC CAC CAC CAC CAC CAC CAC CAT	424
G Y P N E E G H F A S Q H H R G H H H H H H H H H H	76
CAC CAC CAC CAG CAG CAG CAC CAG GCT CTG CAA ACC AAC TGG CAC CTC CCG CAG ATG TCC TCC CCG CCA AGC	499
H H H Q Q Q H Q A L Q S N W H L P Q M S S P P S	101
CGG CCC CGG CAC AGC CTT TGC CTG CAG CCT GAT TCC CGA GGG CCC CCG GAG CTG GGG AGC ACC CCT CCG GTC CTC	574
A A R H S L C L Q P D S G G P P E L G S S P P V L	126
TCC TCC AAC TCT TCT AGC CTG CGC TCC AGC ACC CGG ACC GGA CGC GCG TGC GCA CCA AGG GAT TAT GGC CGT CAA	649
C S N S S S S L G S S T P T G A A C A P R D Y G R Q	151
CGG CTG TCA CCC GCA GAA CTG GAG AAC AGA ACT CGC AGC AAA AGA MA AGC GAC AGT TCA GAT TCC CAG GAA CGA	724
A L S P A E V E K R S G S K R K S D S Q E G	176
ATT TAC AAC TCA GAA GTG AAC AGC AAA CCT AGG AGG GAA AGA ACA CCT TTC ACC AAA GAG CAA ATC AGA GAA CTT	799
N Y K S E V N S X P R K E R T A F T K E Q I R E L	201
GAG GCA GAG TTC GCC CAT CAT AAC TAT CTG ACC AGA CTG AGA AGA TAT GAG ATA CGG GTC AAC CTC GAC CTC ACT	874
Z A E P A H H N Y L T R L R R Y E I A V N L D L T	226
GAA AGA CAG GTG AAA GTG TGG TTC CAG AAC AGG AGA ATG AAG TGG AAG CGG GTC AAG GGG GGA CAA CAA GGA CCT	949
Z R Q V K V N F Q N R R H K W K R V K G G Q Q G A	251
GCA GCC CGA GAA AAG GAA CTG GTG ATT GTG AAA AAG GGA ACA CTT CTT CCA TCA GAG CTG TCA GGA ATT GGT GCA	1024
A A R E K E L V N V K K G T L L P S E L S G I G A	276
GCC ACC CTC CAG CAG ACA CGG GAC TCA CTA GCA ATT GAC GAC AGT CGC GAT AGT GAC CAC AGC TCT GAG CAC GCA	1099
A T L Q Q T G D S L A N D D S R D S D H S S E H A	301
CAC TTA TCA TACATACAGAGACCAGCTCGTCTCAGGAAGCACCATTGTGATGGCAATTCTCACCCAAACATGTTACATGGCAGATGACTGT	1196
H L Stop	303
GCAGTGTGTTAATATAATTAAACGCAAGGCATCTCAAGTGTGTTCTCATGATTGATAGAAGGTTACACTAAGTGCCTCTTATGAAAGATGCTTCCAC	1296
AGTGAATTGGAGAAAGTGAACATATCTAAATATACTTGTGCTTATGACAGAGAGGGAGATGAATGTTGCTTGGCTTGCACTGAAAATTAAATTG	1396
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TGATGAAAATAGGCTTCCCGTGGACAAATGCAGCCAGCTCTGTTATACATTTCGACTCAGCACATCACTATGCTTACTTGTGTT	1796
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TAGAGAACACACCTAGCCCCCTCCAGCCCTCTGTTATGCTGAAAGACATTAGCGTCATAGGCAAGTAGTTACCTTGGCAAATGAGTCTTGTG	1996
CAGATGCTGATTTGTATCTTAAACTGTTAATGGTATGTTGCTGCTTCAGTTAACAGGAAAAGATTCTTCCTCATGTTATGATAACAAACCA	2096
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ATCCAGTACAAGTCCTTGATTTCTGAAAAAGTTAAATGATG	2244

Figure 2

<i>Nmyc-1</i>	■	□	□	■	□	□	■	□	■	
<i>Lamb-1</i>	■	□	■	□	□	□	■	□	■	
<i>Gax</i>	■	□	■	□	■	□	■	□	■	
<i>D12Nyul</i>	■	□	■	□	■	□	■	□	■	
<i>Spnb-1</i>	■	□	■	□	■	□	■	□	■	
	50	57	10	6	4	5	6	4	6	13

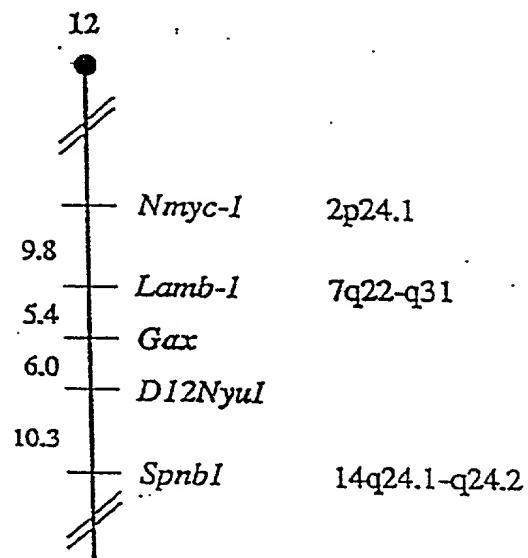


Figure 3

Figure 4

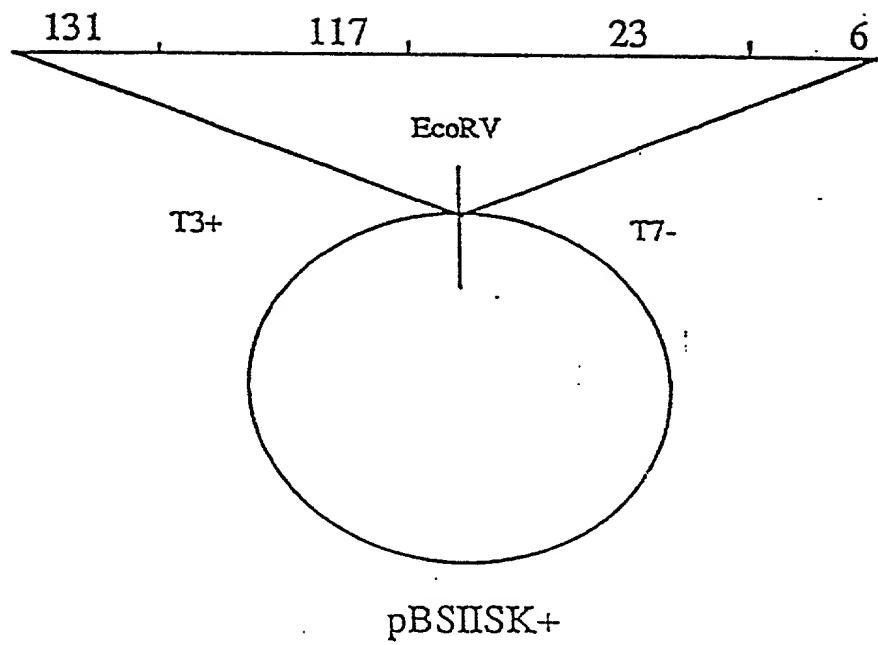


Figure 5a

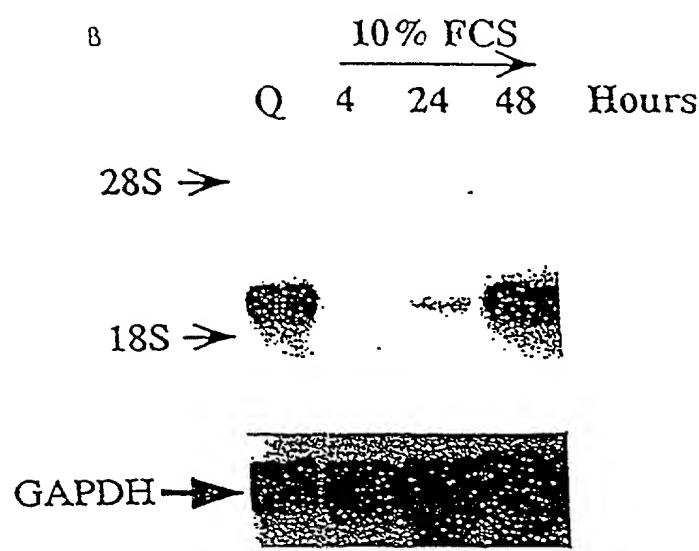


Figure 5B

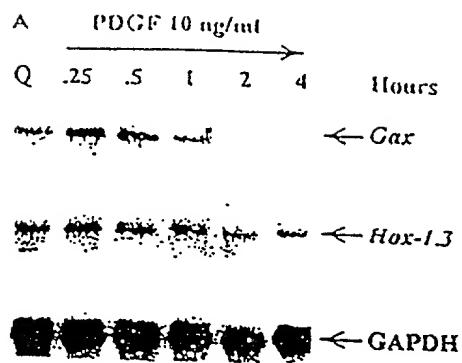
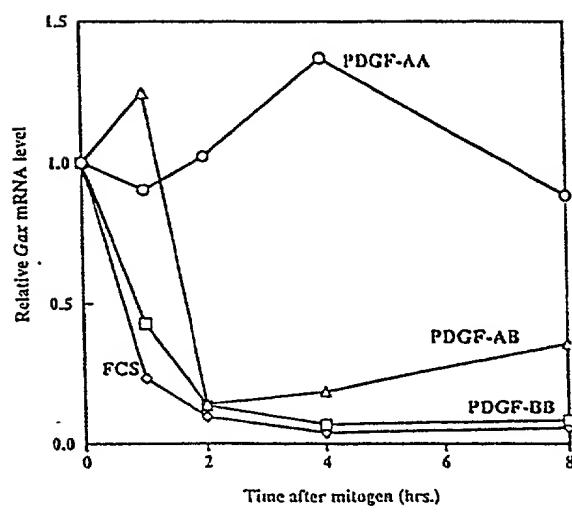


Figure 6



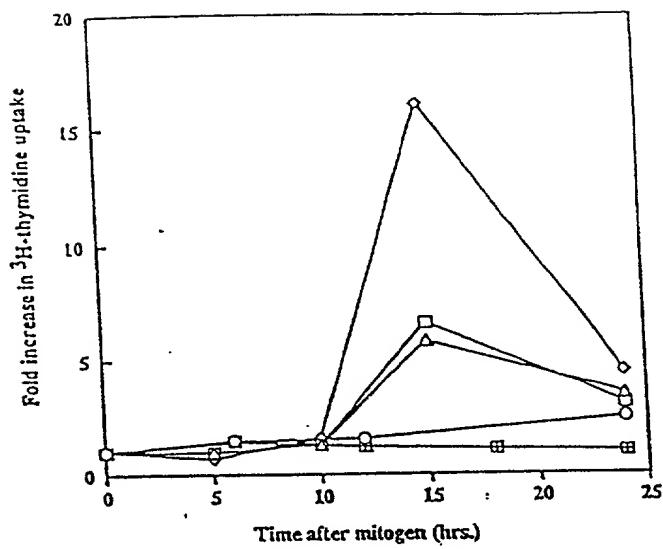


Figure 7

Figure 8

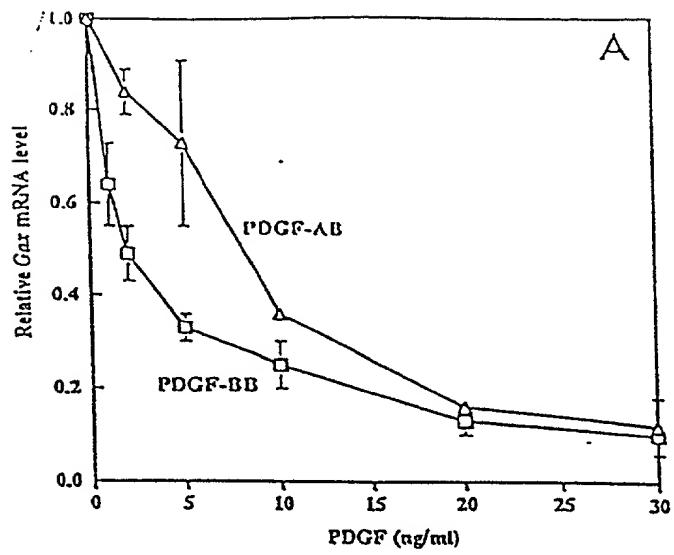


Figure 10

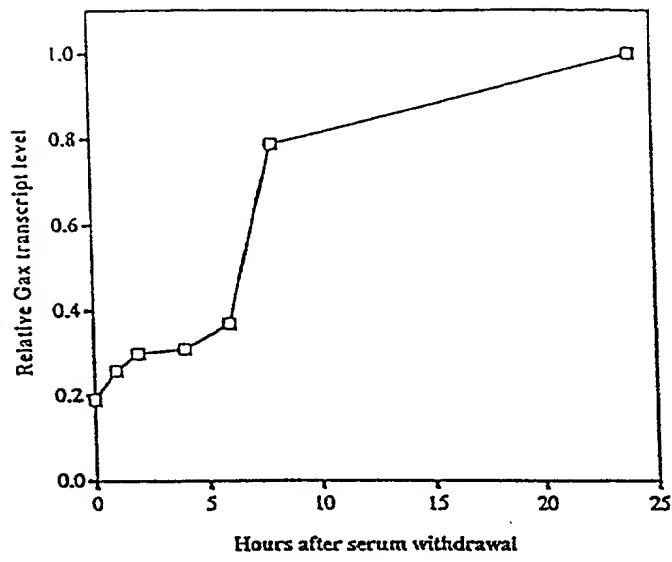


Figure 11

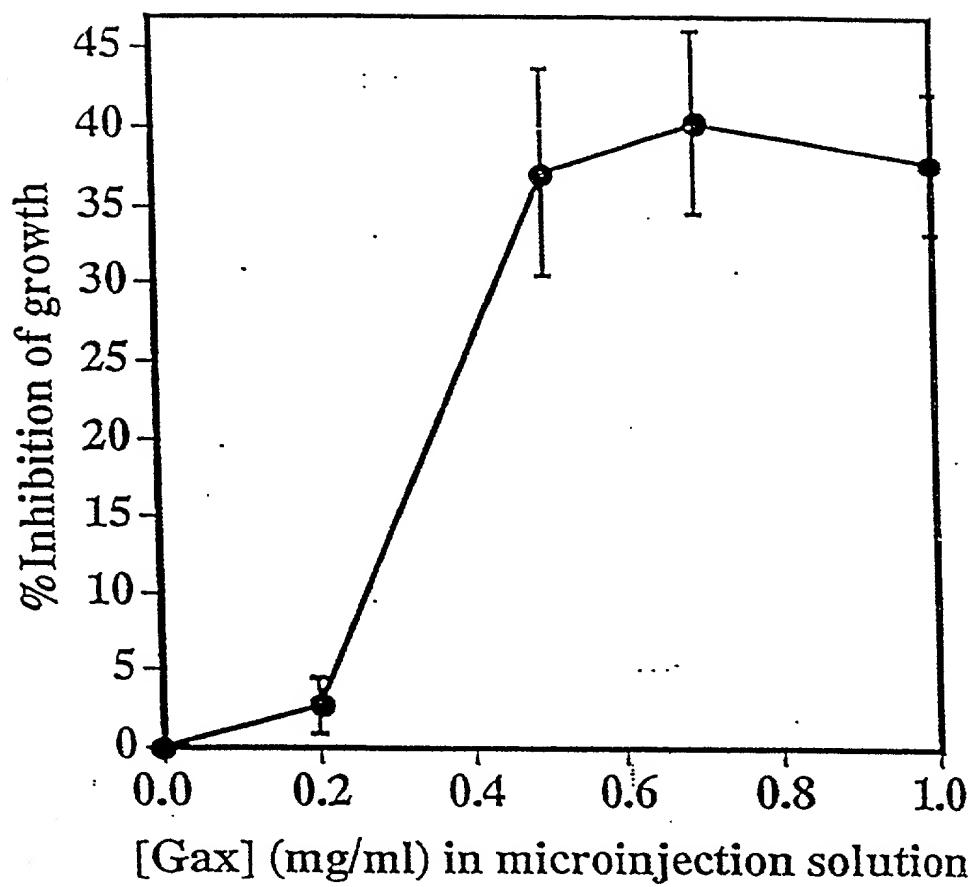


Figure 12

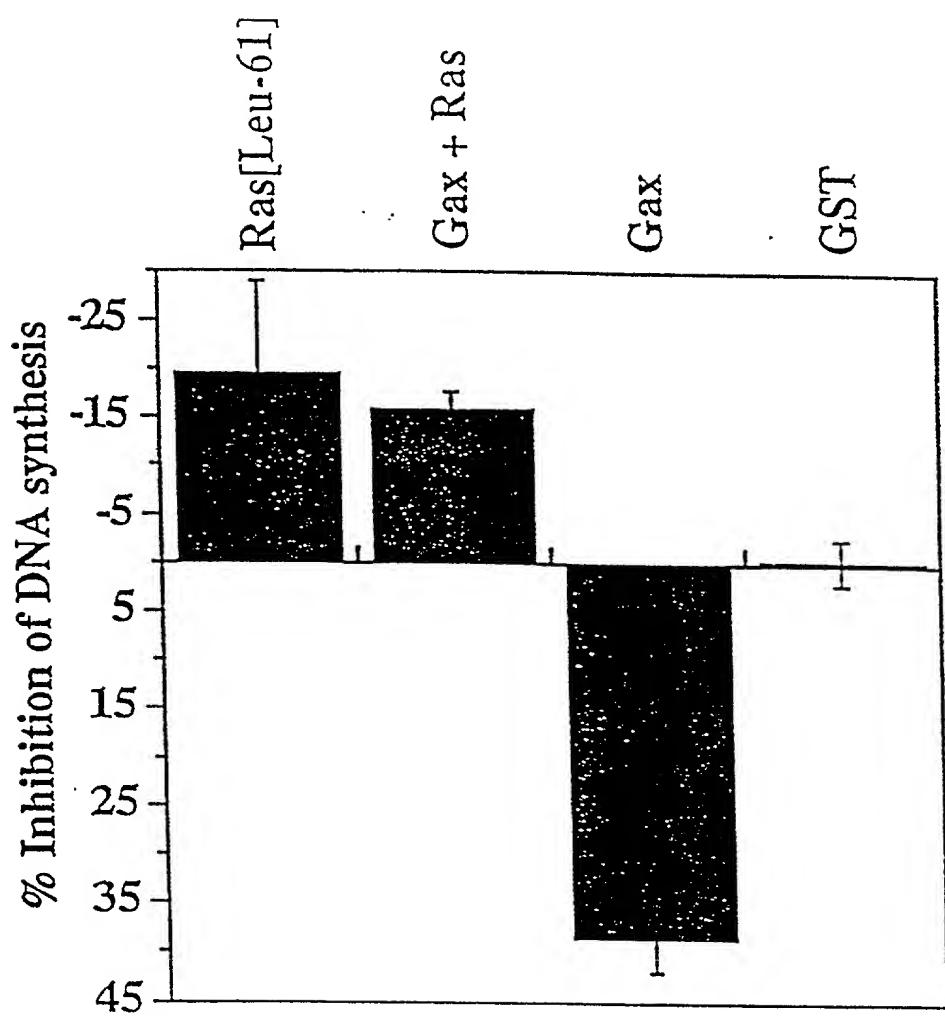


Figure 13B

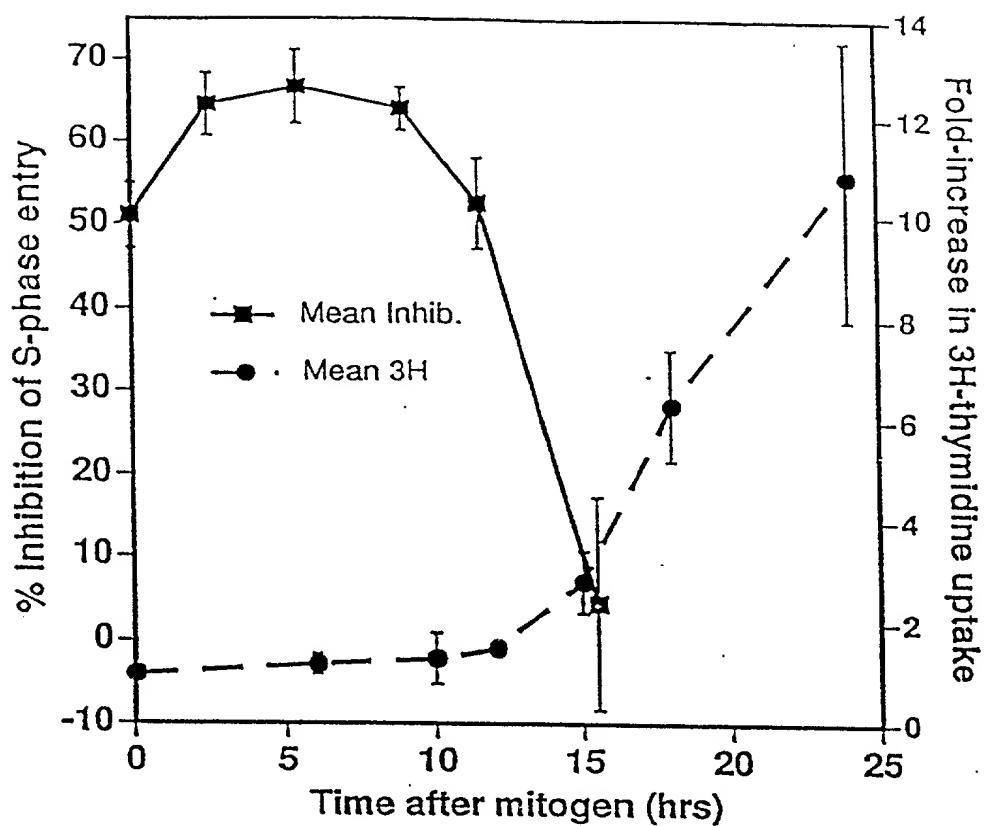


Figure 14

